Dotplot for stranded using anova_p_value HALLMARK_UNFOLDED_PROTEIN_RESPONSE HALLMARK_MYC_TARGETS_V2 HALLMARK_KRAS_SIGNALING_UP HALLMARK XENOBIOTIC METABOLISM HALLMARK MYOGENESIS HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY HALLMARK G2M CHECKPOINT HALLMARK_MYC_TARGETS_V1 HALLMARK MTORC1 SIGNALING HALLMARK TNFA SIGNALING VIA NFKB HALLMARK E2F TARGETS HALLMARK_P53_PATHWAY HALLMARK_IL2_STAT5_SIGNALING HALLMARK_TGF_BETA_SIGNALING HALLMARK_COMPLEMENT HALLMARK_APOPTOSIS HALLMARK_MITOTIC_SPINDLE HALLMARK_INFLAMMATORY_RESPONSE y value HALLMARK_COAGULATION HALLMARK_APICAL_JUNCTION 0.25 HALLMARK_INTERFERON_ALPHA_RESPONSE 0.50 HALLMARK_INTERFERON_GAMMA_RESPONSE 0.75 HALLMARK ESTROGEN RESPONSE LATE HALLMARK ESTROGEN RESPONSE EARLY HALLMARK_GLYCOLYSIS HALLMARK UV RESPONSE UP y value HALLMARK ANDROGEN RESPONSE HALLMARK HEME METABOLISM 0.75 HALLMARK PI3K AKT MTOR SIGNALING HALLMARK HYPOXIA 0.50 HALLMARK IL6 JAK STAT3 SIGNALING 0.25 HALLMARK ANGIOGENESIS HALLMARK EPITHELIAL MESENCHYMAL TRANSITION HALLMARK ALLOGRAFT REJECTION HALLMARK CHOLESTEROL HOMEOSTASIS HALLMARK DNA REPAIR HALLMARK PANCREAS BETA CELLS HALLMARK HEDGEHOG SIGNALING HALLMARK_PROTEIN_SECRETION HALLMARK UV RESPONSE DN HALLMARK APICAL SURFACE HALLMARK WNT BETA CATENIN SIGNALING HALLMARK NOTCH SIGNALING HALLMARK BILE ACID METABOLISM HALLMARK SPERMATOGENESIS HALLMARK ADIPOGENESIS HALLMARK PEROXISOME HALLMARK FATTY ACID METABOLISM HALLMARK KRAS SIGNALING DN HALLMARK OXIDATIVE PHOSPHORYLATION 0.00 0.25 1.00 p_value